

The present invention discloses eight new reduced dimensionality (RD) triple resonance nuclear magnetic resonance (NMR) experiments for measuring chemical shift values of certain nuclei in a protein molecule. The RD 3D <u>HA,CA,(CO),N,HN NMR</u> experiment and the RD 3D <u>H,C,(C-TOCSY-CO),N,HN NMR</u> experiment are designed to yield "sequential" connectivities, while the RD 3D <u>H</u>^{αβ},<u>C</u>^{αβ},CO,HA NMR experiment and the RD 3D <u>H</u>^{αβ},<u>C</u>^{αβ},N,HN NMR experiment provide "intraresidue" connectivities. The RD 3D <u>H,C,C,H-COSY NMR</u> experiment, the RD 3D <u>H,C,C,H-TOCSY NMR</u> experiment, and the RD 2D <u>H,C,H-COSY NMR</u> experiment allow one to obtain assignments for aliphatic and aromatic side chain chemical shifts, while the RD 2D <u>HB,CB,(CG,CD),HD NMR</u> experiment provide information for the aromatic side chain chemical shifts. In addition, a method of conducting suites of RD triple resonance NMR experiments for high-throughput resonance assignment of proteins and identification of the location of secondary structure elements are disclosed.